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APPLICATION NO.	FILING DATE	FIRST NAMED INVENTOR	ATTORNEY DOCKET NO.	CONFIRMATION NO.
09/839,478	04/20/2001	James W. Schumm	016026-9238	4278
23510	7590 06/03/2003			
MICHAEL BEST & FRIEDRICH, LLP ONE SOUTH PINCKNEY STREET P O BOX 1806			EXAMINER	
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MADISON, WI 53701			ART UNIT	PAPER NUMBER
			1634	

DATE MAILED: 06/03/2003

Please find below and/or attached an Office communication concerning this application or proceeding.

	Application No.	Applicant(s)				
Office Action Summary	09/839,478	SCHUMM ET AL.				
cines rioden dummary	Examiner	Art Unit				
The MAILING DATE of this communication app	Jeanine A Goldberg	1634				
Period for Reply	rears on the cover sheet wi	ur the correspondence address				
A SHORTENED STATUTORY PERIOD FOR REPLY THE MAILING DATE OF THIS COMMUNICATION. - Extensions of time may be available under the provisions of 37 CFR 1.13 after SIX (6) MONTHS from the mailing date of this communication. - If the period for reply specified above is less than thirty (30) days, a reply If NO period for reply is specified above, the maximum statutory period version in Failure to reply within the set or extended period for reply will, by statute, any reply received by the Office later than three months after the mailing earned patent term adjustment. See 37 CFR 1.704(b). Status	36(a). In no event, however, may a now within the statutory minimum of thirt will apply and will expire SIX (6) MON a cause the application to become AR	eply be timely filed y (30) days will be considered timely. THS from the mailing date of this communication.				
1) Responsive to communication(s) filed on <u>08 J</u>	lanuary 2003	•				
	is action is non-final.					
3) Since this application is in condition for alloward closed in accordance with the practice under the second sec	ance except for formal mat	ters, prosecution as to the merits is D. 11, 453 O.G. 213				
Disposition of Claims	•					
4) Claim(s) $\underline{22-25}$ is/are pending in the application						
4a) Of the above claim(s) is/are withdrav	vn from consideration.					
5) Claim(s) is/are allowed.						
6) Claim(s) <u>22-25</u> is/are rejected.						
7) Claim(s) is/are objected to.						
8) Claim(s) are subject to restriction and/or	election requirement.					
Application Papers	•					
9) The specification is objected to by the Examiner						
10) ☐ The drawing(s) filed on is/are: a) ☐ accep	_					
Applicant may not request that any objection to the 11) The proposed drawing correction filed on		• •				
If approved, corrected drawings are required in rep		sapproved by the Examiner.				
12) The oath or declaration is objected to by the Exa						
Priority under 35 U.S.C. §§ 119 and 120						
13) Acknowledgment is made of a claim for foreign	priority under 35 H.S.C. 8	119(a)-(d) or (f)				
a) ☐ All b) ☐ Some * c) ☐ None of:	priority under de d.e.c. s	(1).				
1.☐ Certified copies of the priority documents	have been received					
Copies of the certified copies of the priori application from the International Bur See the attached detailed Office action for a list of the certified services.	ity documents have been i eau (PCT Rule 17.2(a)).	received in this National Stage				
14) Acknowledgment is made of a claim for domestic	•					
a) ☐ The translation of the foreign language prov 15)⊠ Acknowledgment is made of a claim for domestic	visional application has be	en received.				
Attachment(s)						
) Notice of References Cited (PTO-892)) Notice of Draftsperson's Patent Drawing Review (PTO-948)) Information Disclosure Statement(s) (PTO-1449) Paper No(s)		ummary (PTO-413) Paper No(s) Iformal Patent Application (PTO-152)				

DETAILED ACTION

- 1. This action is in response to the papers filed January 8, 2003. Currently, claims 22-25 are pending. All arguments have been thoroughly reviewed but are deemed non-persuasive for the reasons which follow. This action is made FINAL.
- 2. Any objections and rejections not reiterated below are hereby <u>withdrawn</u> in view of applicant's arguments, the amendments to the claims and the declaration filed.
- 3. With respect to the inquiry posed to the applicant on March 28, 2003 with respect to a meeting abstract or additional documents from the Fourth International Symposium on Human Identification, the examiner has not received a response. Therefore, the 102(a) has been withdrawn in view of the declaration, however, the examiner has not been able to make a final determination with respect to a 102(b) type inquiry.

Maintained Rejections

Claim Rejections - 35 USC § 103

The following is a quotation of 35 U.S.C. 103(a) which forms the basis for all obviousness rejections set forth in this Office action:

- (a) A patent may not be obtained though the invention is not identically disclosed or described as set forth in section 102 of this title, if the differences between the subject matter sought to be patented and the prior art are such that the subject matter as a whole would have been obvious at the time the invention was made to a person having ordinary skill in the art to which said subject matter pertains. Patentability shall not be negatived by the manner in which the invention was made.
- 4. This application currently names joint inventors. In considering patentability of the claims under 35 U.S.C. 103(a), the examiner presumes that the subject matter of the various claims was commonly owned at the time any inventions covered therein

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were made absent any evidence to the contrary. Applicant is advised of the obligation under 37 CFR 1.56 to point out the inventor and invention dates of each claim that was not commonly owned at the time a later invention was made in order for the examiner to consider the applicability of 35 U.S.C. 103(c) and potential 35 U.S.C. 102(e), (f) or (g) prior art under 35 U.S.C. 103(a).

5. Claims 22-25 are rejected under 35 U.S.C. 103(a) as being unpatentable over Caskey (US. Pat 5,364,759) and Kimpton (Int. J. Leg. Med, 1994) in view of Kimpton (PCR Methods and Applications, 1993) or Fregeau (BioTechniques, 1993) or Urquhart (Int. J. Leg. Med, August 1994).

Caskey discloses the claimed method that includes obtaining a DNA sample, amplifying STR sequences from the DNA sample, and evaluating the amplification products for identification (col. 7, lines 4-10). Caskey teaches a definition of "multiplex polymerase chain reaction (mPCR)"(col. 6, lines 34-68). Further, Caskey teaches that mPCR includes a) primers composed of similar GC base compositions and lengths, b) longer extensions times up to 8 fold the normally utilized times and c) minimization of the number of PCR cycles performed to achieve detection. Caskey teaches that mPCR reaction is optimized for each reaction (col. 6, lines 65-66). Caskey identifies STR loci by searching all human sequences in GenBank (Example 1, col. 8). Strategies to determine the sequences flanking STRs are disclosed in Example 3 (col. 10). Although Caskey teaches that in a reaction with HUMARA and HUMFABP alleles appear as widely spaced doublets such that adjacent alleles overlap, different label may be applied to the different loci to unambiguously identify the alleles (col. 18). Caskey

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teaches the primers of SEQ ID NO: 15, 16, 19, 20, 27, and 28 as primers for the amplification of HUMFABP, HUMPRTB, and HUMTH01, respectively. As described in Example 7, the comparison of amplified alleles by polyacrylamide gel electrophoresis and visualization of the DNA by fluorescent analysis. STR markers can be detected with non-denaturing and denaturing electrophoretic systems (limitations of Claim 29). Silver staining detection methods are all applicable (limitations of Claim 30). Additionally, the loci are selected so that the amplification products of the alleles from different loci do not over lap (limitations of Claim 32). Further, Caskey teaches that the source of DNA to be tested can be any medial or forensic sample and can include blood, semen, vaginal swabs, tissue, hair, saliva, urine and mixtures of body fluids (col. 6, para. 2)(limitations of Claim 33). Caskey discloses the use of allelic ladders as internal standards (col. 7, lines 15-19, and col. 19, lines 15-18)(limitations of Claim 28). Additionally, Caskey teaches kits which contains a container having oligonucleotide primer pair for amplifying STRs and optionally, standards (col. 8 and col. 21, Example 10)(limitations of Claim 34-39, 55).

Kimpton (Int. J. Leg. Med) teaches a multiplex amplification of four tetrameric STR loci. Kimpton (Int. J. Leg. Med) further teaches adjustment of most of the conditions of the multiplex system to optimize results (pg. 303-309). For example, Kimpton (Int. J. Leg. Med) teaches buffer concentration, primer concentration, deoxynucleotide triphosphate concentration, Taq polymerase concentration, template DNA concentration, number of amplification cycles, denaturing temperature, effect of

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annealing temperature, ionic strength and pH, and gel type variation to optimize the multiplex system.

Neither Caskey nor Kimpton specifically teach the combinations of loci recited in the instant claims.

However, Fregeau teaches DNA typing with fluorescently tagged STRs for a sensitive and accurate approach to human identification. Fregeau teaches a multiplex system which contains HUMCD4, HUMFABP, and HUMCATBP2 (pg. 114, col. 3)(limitations of Claim 21, 48-54). DNA for the multiplex was extracted from blood, hair roots, dried bloodstains (pg. 101, col. 3, para. 1)(limitations of Claim 33). Fregeau demonstrates that primers for STR systems HUMHPRT, HUMTH01, HUARA, HUMCD4, HUMFABP, HUMPLA2A1 and HUMRENA4 were used to amplify genomic DNA (pg. 102, col. 1, and Table 1). Fregeau teaches primers identical to the primers of SEQ ID NO: 1, 2, 9, 15, 16, 19, 20, 27, 28, and 30 (Table 1)(limitations of Claim 27). Fregeau teaches HUMvWF, HumFABP, HumACTBP2 and D21S11 all have the same annealing temperature of 64 to 65 degrees and have shown to permit multiplex amplification which saves in reagents and sample template (pg. 117, col. 3, para 2). Further, HumCD4, HumARA, HumTHO01 have the same optimal annealing temperature, 68 degrees. The STR alleles were then separated and detected on a denaturing polyacriamide gel electrophoresis (pg. 106)(limitations of Claim 29). The fluorescent amplification products were resolved on polyacrylamide gels with various gel parameters varied (pg. 103, col. 1)(limitations of Claim 31). A comparison was made between allele sized from silver-stained polyacrylamide gels and automated fluorescent

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analysis (pg. 110, col. 3)(limitations of Claim 30). A four STR system, HUMCD4, HUMHPRT, HUMTH01, HumARA, was explored using additional amplification cycles. Fregeau describes multiplex amplification of polymorphic STR sequences of loci including HUMHPRTB, HUMTH01, HUMCD4, HUMFABP and HUMPLA2A (pg. 117, col. 3, para. 2)(limitations of Claim 26). Empirical evaluation, a specific annealing temperature for each of the STR systems was found to generate consistent allelic profiles with high specificity and sensitivity after 28 cycles of amplification (pg. 115, col. 1). Several benefits of STRs analysis was elucidated including minimal only amounts of template DNA need to be used, the STR alleles can be resolved on sequencing gels using radiolabeled primers or having been processed with cold primers and detected after silver staining, and STRs are amenable to automation (pg. 100-101)(limitations of Claim 30). Further, Fregeau teaches that careful selection of a refined polyacrylamide gel system and appropriate STR loci that have allele size ranges that are mutually resolvable should allow for additional systems to be analyzed with the same fluorescent tag (pg. 117, col. 3).

Kimpton describes the multiplex amplification of polymorphic STR sequences of loci including HUMVWA31, HUMTH01, HUMF13A1, HUMFESFPS, HUMCD4, HUMDHER, HUMCYARO3, HUMAPOAII, HUMPLA2A, HUMIIDA, HUMFABP, HUMGABGA, HUMACTBP2 and D21S11. In Kimpton the combinations of loci are not identical to the combinations claimed. However, Kimpton performs multiplex amplification of STR containing loci in combinations of two, three, four, and seven, chosen loci from HUMVWA31, HUMTH01, HUMF13A1, HUMFESFPS, HUMCD4,

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HUMDHER, HUMCYARO3, HUMAPOAII, HUMPLA2A, HUMIIDA, HUMFABP, HUMGABGA, HUMACTBP2 and D21S11. Kimpton teaches primers for the amplification of HUMACTBP2, HMAPOAII, HUMFABP, HUMTH01, HUMvWA31/A which are identical to the primers taught in the instant application, namely SEQ ID NO: 1, 4, 15, 27, and 32. Kimpton teaches the PCR component concentrations and cycling parameters were optimized for each loci individually. The STRs suitable for co-amplification (multiplexing) were then selected on the basis of similar optimal reaction conditions and compatible allele size ranges (pg. 16, col. 1). Efficient amplification of all loci in multiplex systems was achieved by the adjustment of annealing temperature and individual primer concentration (pg. 19, col. 3). Further, STR loci with overlapping allele size ranges were differentiated by use of different fluorescent dye labels (pg. 16, col. 1).

Urquhart teaches a method of simultaneously determining the alleles present in at least two STR loci. Urquhart teaches a method of preparing DNA from whole blood and performing a PCR amplification using genomic DNA. Each of two primers for each locus were added to the mixture and PCR was performed. The PCR products were electrophoresed in agarose gels, purified and sequenced (pg. 14, col. 1-2)(limitations of Claim 21-22, 26, 48, 50-51). Urquhart teaches primers which are identical to SEQ ID NO: 10, 15, 27, and 32 (Table 1)(limitations of Claim 27). Urquhart also teaches primers which are very homologous to SEQ ID NO: 11, 16, 25, 26 and 31. The alleles were evaluated by separating sizing alleles with an allelic ladder (pg. 14, col. 1)(limitations of Claim 28). Further, Urquhart teaches markers used in the quadruplex STR system were labeled fluorescently (pg. 13-14). The DNA obtained was prepared

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from blood (pg. 14, col. 1)(limitations of Claim 33). The conditions for the reaction were optimized in respect to the different STR's incorporated into the reaction (pg. 14, col. 2). The primers used in the study were all derived from the published or GenBank sequences (pg. 14, col. 1). Although Urquhart does not specifically teach **all** of the recited combinations disclosed in the instant application, Urquhart, does teach the amplification of HUMVWFA31, HUMTH01, HUMF13A01, HUMFES/FPS, HUMCD4, HUMPLA2A1, HUMFOLP23, HUMCYAR04, HUMTFIIDA, HUMFABP, HUMGABRB15, and HUMD21S11 (pg. 14, col. 2). Urquhart teaches that the annealing temperature for HUMTH01, HUMCD4, HUMPLA2A1, HUMFOLP23, HUMCYAR04, HUMTFIIDA, HUMFAB, HUMGABRB15 and HUMD21S11 are all 60 degrees (pg. 14, col. 2).

Therefore, it would have been <u>prima facie</u> obvious to one of ordinary skill in the art at the time the claimed invention was made to have modified the teachings of Caskey and Kimpton (Int. J. Leg. Med) with the loci of Fregeau, Kimpton or Urquhart to obtain the claimed invention based on the teachings of Caskey and Kimpton (Int. J. Leg. Med) in view of Fregeau, Kimpton or Urquhart because the skilled artisan would have been motivated by the teachings of Fregeau, Kimpton, or Urquhart to choose any reasonable number of known STR containing loci, and use them in desired combinations for detection and analysis of polymorphisms in STR loc. Further, it would have been obvious to have chosen any number of known STR containing loci which can be co-amplified together including those suggested by Fregeau and use them in desired combinations for detection and analysis of polymorphisms in STR loci, because such a co-amplification was in fact performed by Kimpton, Fregeau and Urquhart. Both

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Kimpton (Int. J. Leg. Med), Fregeau, Kimpton and Urquhart teach intricate details of multiplex PCR reactions, such as critical parameters for primer design, optimization of cycling conditions, and pros and cons of gel electrophoresis, and visualization techniques (silver stain vs. fluorescence). Both Kimpton and Fregeau references comment on the empirical nature of selecting primers and amplification conditions to achieve an appropriate multiplex amplification system. Kimpton teaches "STRs suitable for co-amplification were selected on the basis of similar optimal reaction conditions and compatible allele size ranges" (pg. 16, col. 1, para 3). For example, Fregeau teaches, HUMTH01 and HUMCD4 both have annealing temperatures of 68 degrees, and have different allele size (bp) which do not overlap (Table 1 and Table 3). Similarly, ACTBP2 and HUMFABP both have annealing temperatures of 64 degrees and do not have overlapping allele sizes (Table 1 and Table 3). Therefore the at least two STR loci would contain clearly distinguishable STR allelic profiles (pg. 115, col. 3) and would have been obvious to combine the two STR loci to obtain the claimed invention. The choice of STR loci chosen to multiplex is dependent on what information is desired from the allele analysis. As exemplified in the art, gel analysis of several STR loci on the same gel saved time and reagents. One of ordinary skill in the art would have been motivated to design appropriate primers and optimize PCR conditions in order to co-amplify additional combinations of STR loci for the benefit of saving time, reagents and other supplies in the amplification process as taught by Fregeau (pg. 117). It is also evident from these references that the loci were chosen for their already demonstrated polymorphic properties and that implementation of multiplex amplification

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of combinations thereof was easily achieved by routine optimization of the well known PCR methodology adapted for multiplex purposes. As admitted in the specification, "successful combinations are generated by trial and error (routine experimentation) of locus combinations and by adjustment of primer concentrations to identify an equilibrium in which all included loci may be amplified"(pg. 10, lines 10-13). Thus, the claimed invention would have been obvious over Caskey and Kimpton (Int. J. Leg. Med) in view of Fregeau, Kimpton or Urquhart.

Response to Arguments

The response traverses the rejection. The response asserts that the references indicate that the selection of STR loci that can be co-amplified is not a trivial matter, but rather one that would require a considerable amount of experimentation. This argument has been reviewed but is not convincing because the standard for obvious is not absolute expectation of success, but rather reasonable expectation of success. Given the teachings of the references there is a reasonable expectation of success. While some routine experimentation and optimization may be required to determine the exact parameters which allow successful optimization of the assay, this routine optimization is not an indice of non-obviousness. As noted in *In re Aller*, 105 USPQ 233 at 235, "More particularly, where the general conditions of a claim are disclosed in the prior art, it is not inventive to discover the optimum or workable ranges by routine experimentation." Routine optimization is not considered inventive and no evidence has been presented that the probe selection performed was other than routine, that the products resulting from the optimization have any unexpected properties, or that the results should be

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considered unexpected in any way as compared to the closest prior art. It is noted that this is not an invitation to file a declaration after final. As provided by MPEP 716.01, "Evidence traversing rejections must be timely or seasonably filed to be entered and entitled to consideration. In re Rothermel, 276 F.2d 393, 125 USPQ 328 (CCPA 1960). Affidavits and declarations submitted under 37 CFR 1.132 and other evidence traversing rejections are considered timely if submitted:(1) prior to a final rejection...."

MPEP 716.01(c) makes clear that "The arguments of counsel cannot take the place of evidence in the record. In re Schulze, 346 F.2d 600, 602, 145 USPQ 716, 718 (CCPA 1965). Examples of attorney statements which are not evidence and which must be supported by an appropriate affidavit or declaration include statements regarding unexpected results, commercial success, solution of a long - felt need, inoperability of the prior art, invention before the date of the reference, and allegations that the author(s) of the prior art derived the disclosed subject matter from the applicant." Here, the statements regarding the inoperability of the prior art must be supported by evidence, not argument.

The response disagrees with the Examiner's characterization of Urquhart because the response asserts that only four STR loci were suitable candidates for multiplex amplification and analysis. This argument has been thoroughly reviewed, but is not found persusasive because the references does not state that the STR loci not included in the multiplex analysis were unsuitable. The references states that "major considerations for selection of loci were discriminating power, absence of linkage, agreement with Hardy-Weinberg equilibrium, low levels of shadow bands, compatibility

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with other loci and accurate sizing of alleles. This passage does not indicate that the other STR loci were unsuitable candidates, but merely provides a single example of a combination of loci for multiplex. Thus for the reasons above and those already of record, the rejection is maintained.

1. Claims 22-25 are rejected under 35 U.S.C. 103(a) as being unpatentable over Caskey (5,364,759) in view of GenBank STR lociHUMTH01, HUMTPOX, HUMF13A01, HUMFABP, HUMMYPOK, HUMBFXIII, HUMHPRTB, HSAC04, HUMCYP19 and HUMPLA2A1.

Caskey discloses the claimed method that includes obtaining a DNA sample, amplifying STR sequences from the DNA sample, and evaluating the amplification products for identification (col. 7, lines 4-10). Caskey describes a preferred Caskey discloses the claimed method that includes obtaining a DNA sample, amplifying STR sequences from the DNA sample, and evaluating the amplification products for identification (col. 7, lines 4-10). Caskey teaches a definition of "multiplex polymerase chain reaction (mPCR)"(col. 6, lines 34-68). Further, Caskey teaches that mPCR includes a) primers composed of similar GC base compositions and lengths, b) longer extensions times up to 8 fold the normally utilized times and c) minimization of the number of PCR cycles performed to achieve detection. Caskey teaches that mPCR reaction is optimized for each reaction (col. 6, lines 65-66). Caskey identifies STR loci by searching all human sequences in GenBank (Example 1, col. 8). Strategies to determine the sequences flanking STRs are disclosed in Example 3 (col. 10). Although

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Caskey teaches that in a reaction with HUMARA and HUMFABP alleles appear as widely spaced doublets such that adjacent alleles overlap, different label may be applied to the different loci to unambiguously identify the alleles (col. 18). Caskey teaches the primers of SEQ ID NO: 15, 16, 19, 20, 27, and 28 as primers for the amplification of HUMFABP, HUMPRTB, and HUMTH01, respectively. As described in Example 7, the comparison of amplified alleles by polyacrylamide gel electrophoresis and visualization of the DNA by fluorescent analysis. STR markers can be detected with non-denaturing and denaturing electrophoretic systems (limitations of Claim 29). Silver staining detection methods are all applicable (limitations of Claim 30). Additionally, the loci are selected so that the amplification products of the alleles from different loci do not over lap (limitations of Claim 32). Further, Caskey teaches that the source of DNA to be tested can be any medial or forensic sample and can include blood, semen, vaginal swabs, tissue, hair, saliva, urine and mixtures of body fluids (col. 6, para. 2)(limitations of Claim 33). Caskey discloses the use of allelic ladders as internal standards (col. 7, lines 15-19, and col. 19, lines 15-18)(limitations of Claim 28). Additionally, Caskey teaches kits which contains a container having oligonucleotide primer pair for amplifying STRs and optionally, standards (col. 8 and col. 21, Example 10)(limitations of Claim 34-39, 55).

Caskey does not specifically teach the recited locus combinations.

However, the STR loci HUMTH01, HUMTPOX, HUMF13A01, HUMFABP, HUMMYPOK, HUMBFXIII, HUMHPRTB, HSAC04, HUMCYP19 and HUMPLA2A1 have been taught by GenBank Accession No: HSAC04, M28420, M21986 J03834, M64554

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J05294, M18079 J03465, M26434, M87312, M22970 M14965, D00269, M68651, and M25858 M25716.

Furthermore, rather than citing STR containing loci, Caskey refers to STR sequences by their alphabetical designation as indicated in Table I. Additionally, Caskey does not recite locus combinations in examples 4-7 and tables 6-9, where data from multiplex amplification of said alleles is performed and analyzed. Caskey describes the level of skill of an ordinary artisan by stating that once STR sequences and their flanking sequences are obtained, primer pairs may be designed and synthesized according to the flanking sequences and PCR amplification and comparison of amplified products may be performed to detect the short tandem repeats (col. 4, lines 9-17, col. 5, lines 16-53, col. 6, lines 58-60). Identical primers were used in the instant application for HUMFABP, HUMTH01, and HUMPRTB, therefore, the method by which Caskey derives primers for STR loci appears to be consistent with the method of the instant application. Caskey also comments on the empirical nature of multiplex amplification reactions and points out that each reaction must be optimized (col. 6, line 65).

Therefore, to one of ordinary skill in the art at the time the invention was made, it would have been **prima facie** obvious to use any number of primers, including SEQ ID NO: 1-32, among other possible sequences that could accomplish the same goal for the process of simultaneously amplifying specified loci which provide a different pattern and thus a means of confirmation or subsequent analysis. SEQ ID NO:s 1-14, 17-18, 21-26, 29-32 are not specifically taught by Caskey as specific primers for the respective STR

loci. The claimed primers, however, would have been obvious based on the teaching of Caskey about primer design and synthesis and the known sequences of the claimed loci, which were available from GenBank. Additionally, Caskey was able to perform multiplex amplification of HUMTH01 in combination with other loci, which reiterates the level of skill in the art. As admitted in the specification, "successful combinations are generated by trial and error (routine experimentation) of locus combinations and by adjustment of primer concentrations to identify an equilibrium in which all included loci may be amplified"(pg. 10, lines 10-13). Therefore, the claimed invention would have been obvious over Caskey in view of the GenBank entries.

Response to Arguments

The response traverses the rejection. The response asserts that Caskey provides no teaching as to which loci could be amplified to produce results that could be evaluated in any meaningful way because of the overlapping alleles. This argument has been reviewed but is not convincing because Caskey provides methods for choosing primers for use in multiplex analysis. Further the specification, "successful combinations are generated by trial and error (routine experimentation) of locus combinations and by adjustment of primer concentrations to identify an equilibrium in which all included loci may be amplified"(pg. 10, lines 10-13). Thus for the reasons above and those already of record, the rejection is maintained.

Double Patenting

The nonstatutory double patenting rejection is based on a judicially created doctrine grounded in public policy (a policy reflected in the statute) so as to prevent the unjustified or improper timewise extension of the "right to exclude" granted by a patent and to prevent possible harassment by multiple assignees. See *In re Goodman*, 11 F.3d 1046, 29 USPQ2d 2010 (Fed. Cir. 1993); *In re Longi*, 759 F.2d 887, 225 USPQ 645 (Fed. Cir. 1985); *In re Van Ornum*, 686 F.2d 937, 214 USPQ 761 (CCPA 1982); *In re Vogel*, 422 F.2d 438, 164 USPQ 619 (CCPA 1970);and, *In re Thorington*, 418 F.2d 528, 163 USPQ 644 (CCPA 1969).

A timely filed terminal disclaimer in compliance with 37 CFR 1.321(c) may be used to overcome an actual or provisional rejection based on a nonstatutory double patenting ground provided the conflicting application or patent is shown to be commonly owned with this application. See 37 CFR 1.130(b).

Effective January 1, 1994, a registered attorney or agent of record may sign a terminal disclaimer. A terminal disclaimer signed by the assignee must fully comply with 37 CFR 3.73(b).

6. Claim 22 is rejected under the judicially created doctrine of obviousness-type double patenting as being unpatentable over claims 1-43 of U.S. Patent No. 6,221,598. An obvious –type double patenting rejection is appropriate where the conflicting claims are not identical, but an examined application claim not patentably distinct from the reference claim(s) because the examined claim is either anticipated by, or would have been obvious over, the reference claims(s). See, e.g., In re Berg, 140 F.3d 1428, 46 USPQ2d 1226 (Fed. Cir. 1998); In re Goodman, 11 F.3d 1046, 29 USPQ2d 2010 (Fed. Cir. 1993); In re Longi, 759 F.2d 887, 225 USPQ 645 (Fed. Cir. 1985). Although the conflicting claims are not identical, they are not patentably distinct from each other because claims 21, 48 are generic to all that is recited in claim 1 of U.S. Patent No. 6,221,598. Specifically, the method of the patent in one embodiment is directed to detecting HUMPOX, HUMCD4 and HUMTH01. For example, the instant claims are drawn to methods of simultaneously determining at least two makers selected from

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HUMCD4 and HUMTH01. Therefore, the claimed methods fall within the scope of the claims already patented. The method of Claim 22 in the instant application and the method of Claim 1 of 6,221,598 encompass the same scope. Specifically, the first set of short tandem repeats is the same as those listed in Claim 1 of 6,221,598.

Response to Arguments

The response traverses the rejection. The response asserts that a terminal disclaim for US Pat. 6,221,598 has been filed. This argument has been reviewed but is not convincing because the terminal disclaimer which was filed is not proper. The application/patent being disclaimed has not been identified. The instant application number has not been identified within the body of the terminal disclaimer. Thus for the reasons above and those already of record, the rejection is maintained.

7. Claims 22-23, 25 are rejected under the judicially created doctrine of obviousness-type double patenting as being unpatentable over claims 1-43 of U.S. Patent No. 5843660. An obvious –type double patenting rejection is appropriate where the conflicting claims are not identical, but an examined application claim not patentably distinct from the reference claim(s) because the examined claim is either anticipated by, or would have been obvious over, the reference claims(s). See, e.g., In re Berg, 140 F.3d 1428, 46 USPQ2d 1226 (Fed. Cir. 1998); In re Goodman, 11 F.3d 1046, 29 USPQ2d 2010 (Fed. Cir. 1993); In re Longi, 759 F.2d 887, 225 USPQ 645 (Fed. Cir. 1985). Although the conflicting claims are not identical, they are not patentably distinct from each other because claims 21, 48 are generic to all that is recited in claim 1 of U.S.

Patent No. 6,221,598. Specifically, the method of the patent in one embodiment is directed to detecting HUMCSF1PO, HUMPOX, HUMCD4 and HUMTH01 (limitations of Claim 24) or HUMCSF1PO, HUMPOX, HUMF13A01 and HUMTH01 (limitations of Claim 25). For example, the instant claims are drawn to methods of simultaneously determining at least two makers selected from HUMCD4 and HUMTH01 (limitations of Claim 21); HUMCSF1PO, HUMPOX, and HUMTH01 (limitations of Claim 22). Therefore, the claimed methods fall within the scope of the claims already patented. Claim 4 of the patent is directed to determining a set of seven loci which comprises HUMTPOX, HUMTHO01 and HUMCSF1PO (limitations of Claim 23). Claim 34 of the instant application is drawn to a kit comprising primers for at least two loci which in one embodiment is directed to HUMCSF1PO and HUMTPOX. Claim 25 of 5,743,660 teaches a kit comprising primers from D16S539, D7S820, D13S317, D5S808, HUMCSF1PO, HUMTPOX which encompasses Claim 34. Moreover, Claim 25 of 5,743,660 teaches a kit comprising primers from D16S539, D7S820, D13S317, D5S808, HUMCSF1PO, HUMTPOX, HUMTH01 and HUMvWFA31 which encompasses Claim 36-37 of the instant application. Claim 4 of 5,743,660 teaches a method using D16S539, D7S820, D13S317, D5S818, HUMCSF1PO, HUMTPOX and HUMTH01 (limitations of Claim 40, 41).

Response to Arguments

The response fails to address the rejection. Thus for the reasons above and those already of record, the rejection is maintained.

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8. Claim 22-23, 25 are rejected under the judicially created doctrine of obviousness-type double patenting as being unpatentable over claims 1-4, 6, 9-18, 21-23, 60-36, 39 of copending Application No. 09/199,542, now Patent 6,479,235. Although the conflicting claims are not identical, they are not patentably distinct from each other. The instant claims are drawn to detecting two or more STR loci. The claims of the allowed application are directed to detection of thirteen STR loci. The set of 13 STR loci encompasses the set of 3 loci of the instant application.

This is no longer a <u>provisional</u> obviousness-type double patenting rejection because the conflicting claims have not in fact been patented. It is noted that the claims for 09/199,542 have been allowed and have matured into a patent.

Response to Arguments

The response fails to address the rejection. Thus for the reasons above and those already of record, the rejection is maintained.

Conclusion

9. No claims allowable.

10. Applicant's amendment necessitated the new ground(s) of rejection presented in this Office action. Accordingly, **THIS ACTION IS MADE FINAL**. See MPEP § 706.07(a). Applicant is reminded of the extension of time policy as set forth in 37 CFR 1.136(a).

A shortened statutory period for reply to this final action is set to expire THREE MONTHS from the mailing date of this action. In the event a first reply is filed within

TWO MONTHS of the mailing date of this final action and the advisory action is not mailed until after the end of the THREE-MONTH shortened statutory period, then the shortened statutory period will expire on the date the advisory action is mailed, and any extension fee pursuant to 37 CFR 1.136(a) will be calculated from the mailing date of the advisory action. In no event, however, will the statutory period for reply expire later than SIX MONTHS from the date of this final action.

11. Any inquiry concerning this communication or earlier communications from the examiner should be directed to examiner Jeanine Goldberg whose telephone number is (703) 306-5817. The examiner can normally be reached Monday-Friday from 8:00 a.m. to 5:30 p.m.

If attempts to reach the examiner by telephone are unsuccessful, the examiner's supervisor, Gary Jones, can be reached on (703) 308-1152. The fax number for this Group is (703) 305- 3014.

Any inquiry of a general nature should be directed to the Group receptionist whose telephone number is (703) 308-0196.

Jeanine Goldberg May 20, 2003

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